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UNIVERSITY OF UTAH
SCHOOL OF MEDICINE

Department of Pathology



ACCE and Genomic Testing

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Scope

- Focus on molecular assays, attempting to fit models/definitions developed for clinical chemistry
- Disclaimer: My own opinions, considering BCBSA policy, but also looking towards the future

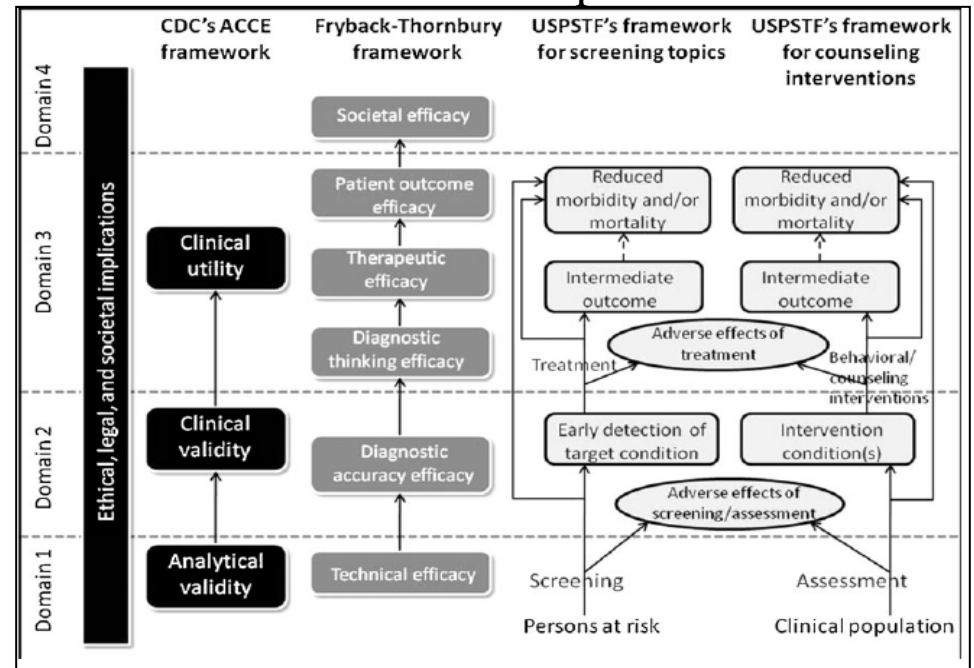
ACCE Framework

- **A**nalytical validity:
- **C**linical Validity
- **C**linical Utility
- **E**thical, legal, social implications
- Purposes for tests

– Reduce morbidity/mortality

- Provide information to manage patient/family members
- Assist with reproductive decision-making

Framework Comparisons



Fryback GDG, Thornbury JR. The efficacy of diagnostic imaging. *Med Decis Making* 1991 11: 88
 Matchar Chapter 1. Introduction to methods guide for medical test reviews. *JGenIntmed* 2012

BioMarkers vs Mutations

- Molecular Biomarkers (associated, relative risks)
 - Clinical trials to establish association
 - More likely to be proprietary
 - GWAS studies
 - Expression patterns
- Pathogenic variants (causative)
 - Mendelian disorders (germline)
 - Oncology (somatic variants)
 - Driver mutations, therapy – drug susceptibility, resistance variants

Analytical Validity

- Does the assay detect what is claimed that it detects?
 - Accuracy/ precision studies
 - Determines analytical sensitivity and specificity
 - Region interrogated defined
 - Targeted mutations
 - Gene sequencing
 - Targeted exons, Full gene sequencing (all exons, intron/exon boundaries, some known deep intronic or regulatory mutations)
 - Deletion/duplication analysis
 - Performance affected by
 - Interfering substances (well known)
 - Rare, unknown variants at primer/probe sites, creating 2^o structure
 - Mosaicism, low mutation levels, limits of detection
 - Continuing evaluation: proficiency testing/alternative assessment

Clinical Validity

- Does the test correctly identify affected/unaffected individuals?
 - Does Analyte (gene) or Assay determine clinical validity?
 - Do mutations in a gene cause disease?
 - Linkage studies, functional analysis, case/control, cloned from known protein sequence
 - Depends on the region interrogated /defined phenotype
 - Clinical sensitivities (F8 example)
 - Not necessarily method dependent
 - PPV/NPV a measure of analytic or clinical validity or clinical utility?
 - How is it defined for single gene disorders?
 - Penetrance, mild vs severe mutations?
 - Dependent on population, indication for testing

Clinical Validity - Complications

- Inherited disease concepts
 - **Penetrance/expressivity**
 - Pleiotropy – single gene influences multiple traits
 - **Clinical Overlap:** pathogenic variants in multiple genes cause similar phenotypes
 - **Phenocopy** – phenotype overlap due to environment that resembles the effect of inherited pathogenic variants
 - Carefully define “phenotype”, (BRCA Example)
 - Polygenic traits: multiple genes contribute to the phenotype
 - **Same test for diagnostic, predictive, carrier testing**
 - Interrogating regions (deep intronic, regulatory) of a gene or genes not well understood will produce more Variants of Uncertain Significance (VUS)
 - All genes on a panel to have established clinical validity
 - ClinGen project funded by NIH to examine disease categories

Modified ACCE (Fryback-Thornbury) for Clinical Utility

- Diagnostic Thinking Efficacy (Diagnosis):
 - Rule out disease (differential diagnosis)
 - Stop diagnostic odyssey: prevent additional testing
 - Appropriate follow-up/monitoring
- Therapeutic efficacy
 - Drug response
- Patient outcome efficacy
 - Patient management: improve outcomes
 - Prognostic: Determine aggressiveness of disease/treatment
 - Predictive: pre-symptomatic, familial mutations, reproductive
- Societal efficacy:
 - Proper use of medical/community resources

Reasons to Show Utility

- Aid clinicians in ordering, interpreting
- Demonstrate value of genomic medicine
- Reimbursement

Definition of Clinical Utility

- Utility for patient, clinician, payers, regulators, society
- Definition of Clinical Utility
 - Narrow: Determine drug and dose – improved outcomes demonstrated
 - For clinician/patient: diagnosis, treatment, management
 - Inherent utility of diagnostic testing
 - For patient/family: predictive testing, reproductive planning, long term care planning
 - For Payers: treatment, improved outcomes
 - For Regulators: analytical and clinical validity, expand to utility?
 - For Society: Efficient use of healthcare/community resources

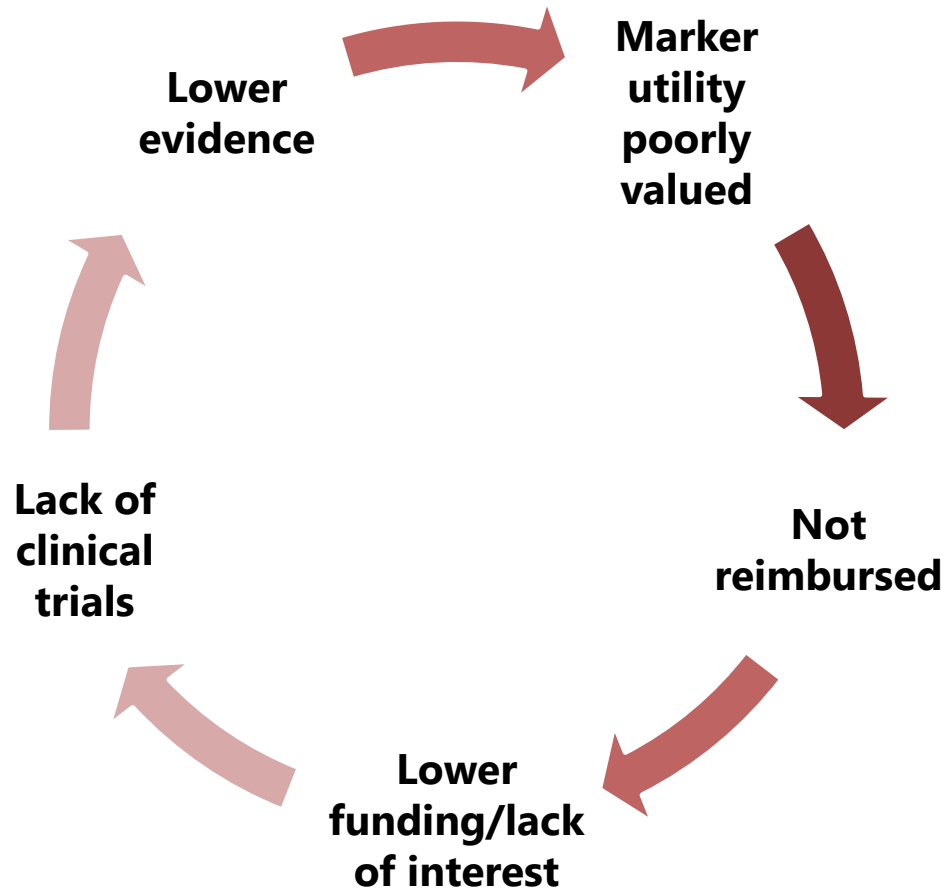
Establishing Clinical Utility

- Randomized prospective controlled studies
- Retrospective studies
 - Archived samples
- Issues with:
 - Rare inherited diseases
 - Rare mutations (somatic)
 - Long duration
 - Ethically valid?
 - Inconclusive results
 - Poorly designed
 - Insufficient numbers

EGAPP

- **E**valuation of **G**enomic **A**pplications in **P**ractice and **P**revention
- Common conclusion
 - Insufficient evidence
 - ...found insufficient evidence to support a recommendation for or against use of CYP450 testing in adults beginning SSRI treatment for non-psychotic depression.
 - In the absence of supporting evidence..., EGAPP discourages the use of CYP450 testing for patients beginning SSRI treatment until further clinical trials are completed
- Taken as “Evidence Against”
 - SSRI studies extended to other uses
- Re-evaluate with continuing studies

Circular Problem



Adapted from: Generating Evidence for Genomic Diagnostic Test Development: Workshop Summary: National Academy of Sciences.
http://www.nap.edu/catalog.php?record_id=13133

Testing Symptomatic Individuals

- Diagnostic:
 - Explain the clinical symptoms
 - Understand disease course
- Prognostic:
 - Understand likely disease progression
 - Preventive management
- Therapeutic:
 - Determine most effective treatment/management

Asymptomatic Individual

- Predictive testing
- Family history
- Known familial mutations
 - Test affected individual for the benefit of family members
- Population screening
 - Newborn screening (State programs)

Testing Cancer Cells (Somatic)

- Diagnostic: identify genetic abnormalities causative of or resulting from disease
- Prognostic: determine aggressiveness of disease/treatment
- Predictive: determine therapy, resistance to therapy

Models

- Fully powered clinical studies not always feasible
 - Underpowered or partial data modeled for useful information?
- Require models for different scenarios? Types
 - Oncology
 - Chain of evidences (biological relationships/pathways)
 - Demonstrative usefulness in one or multiple cancer/specimen types?
 - Define “supportive” and “adequate” evidence
 - Inherited diseases
 - Approximately 4600 known medically relevant genes
 - Show each disease separately?
 - Another 20,000 in genome – how many will be shown to be medically relevant?
 - Compare to non-molecular diagnostic pathway/procedures
 - Diagnostic efficacy
- Same assay used for different purposes

Clinical Utility for Oncology

- “Driver” mutations essential for tumor progression
- “Passenger” mutations that might facilitate, but not essential for progression
- Prognosis
 - Help determine aggressiveness of treatment
- Predictive testing for therapy
 - Multiple tumor types – *BRAF* V600E
 - Histologically identical tumors- *KRAS2*

Selected Molecular Tests with Tier 1 cpt Oncology

CONDITION	DIAGNOSIS	MANAGEMENT	PROGNOSIS	PREDICTIVE
Acute myeloid leukemia	✓	✓	✓	✓
Stem cell transplant monitoring		✓	✓	✓
Chronic lymphocytic leukemia		✓	✓	✓
Chronic myelogenous leukemia	✓	✓	✓	✓
Colon Cancer		✓	✓	✓
Breast and ovarian cancer		✓	✓	✓
Non-small cell lung cancer		✓	✓	✓
Acute promyelocytic leukemia t(15;17)		✓	✓	✓
Gastrointestinal Stromal Tumors		✓	✓	✓
Melanoma		✓	✓	✓

Clinical Utility for Inherited Diseases

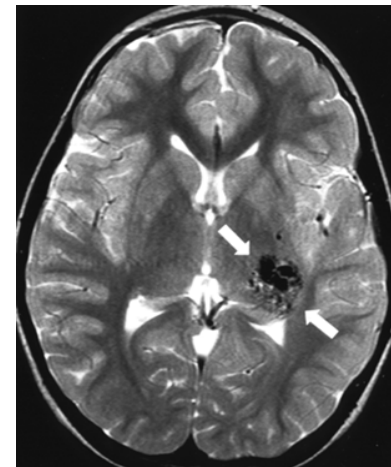
- Many are rare:
 - Approximately 4600 known human genetic disorders
 - Not feasible to show utility for each one
 - Aggregate by disease type, test method?
 - Still may have strong clinical validity/utility
 - lack cpt codes
 - Together, they are substantive
 - 100% of individuals have genetic variants that could affect drug response
 - *JAMA 286:2270, 2001.*

Selected Molecular Tests with Tier 1 cpt Genetics

CONDITION	DIAGNOSIS	MANAGEMENT	PROGNOSIS	PREDICTIVE
Alpha thalassemia, Hb Bart hydrops fetalis syndrome, HbH disease	✓	✓		✓
Alpha-1-antitrypsin deficiency	✓	✓	✓	✓
Ashkenazi Jewish:				
Bloom syndrome	✓	✓		✓
Canavan disease	✓	✓		✓
Tay-Sachs disease	✓	✓	✓	✓
Cardiomyopathies	✓		✓	✓
Cystic Fibrosis	✓			✓
Cytogenomic constitutional abnormalities (e.g. Klinefelter, trisomy 21)	✓			✓
Familial adenomatous polyposis (FAP)	✓		✓	✓
Fragile X	✓		✓	✓
Huntington Disease	✓		✓	✓
Hereditary breast and ovarian cancer	✓		✓	✓
Hereditary hemochromatosis	✓			Limited
Hereditary non-polyposis colorectal cancer, Lynch syndrome	✓		✓	✓
Long QT syndrome	✓		✓	✓
Marfan syndrome	✓	✓	✓	✓
Nonsyndromic hearing loss	✓	✓		✓
Rett syndrome	✓	✓		✓
Spinal Muscular Atrophy	✓	✓	✓	✓

Example: Hereditary Hemorrhagic Telangiectasia

- Appropriate use of health resources
 - Life threatening cerebral/pulmonary manifestations
 - Brain MRI with contrast:
 - Contrast echocardiogram:
 - 20% need F/U of chest CT, radiation exposure
 - Surveillance: every 5 years in affected individuals, or in unaffected individuals until approximately age 40 (unless ruled out by molecular testing)
 - Guidelines available
 - Faughnan J Med Genet 2011;48:73e87



Pictures courtesy of Whitney Wooderchak-Donahue

Single Gene vs Gene Panel

- ASHG:
 - “..., the scope of genetic testing should be limited to single-gene analysis or targeted gene panels based on the clinical presentation of the patient...”
 - Botkin JR et al. Points to consider: Ethical, legal, and psychosocial implications of genetic testing in children and adolescents. ASHG 2015;97:6-21
- Use most focused assay available (as appropriate)
 - Single gene, if meets clinical criteria
 - Small gene panel improves diagnostic yield, if non-classic phenotype
 - Large gene panels - common symptoms for numerous diseases, in place of an exome?
 - Exome/genome for combination of symptoms/family history consistent with genetic etiology, but remains undiagnosed

Marfan syndrome

- Tall stature
- Arachnodactyly
- Hypermobile joints
- Scoliosis
- Aortic aneurysm
- Learning disability
- Positive family history, sudden death in a close relative



http://www.healthinplainenglish.com/health/cardiovascular/marfan_syndrome/index.htm

Loeys-Dietz Syndrome

- Arterial tortuosity
- Hypertelorism
- Bifid (split) or broad uvula
- Aneurysms
- Scoliosis
- Positive family history, sudden death in a close relative



McGee et al., 2011, Circulation

Ehler Danlos Syndrome Type IV

- Aneurysm
- Thin, translucent skin
- Extensive bruising
- Hypermobility
- Clubfoot
- Spontaneous pneumothorax or haemothorax
- Positive family history, sudden death in a close relative



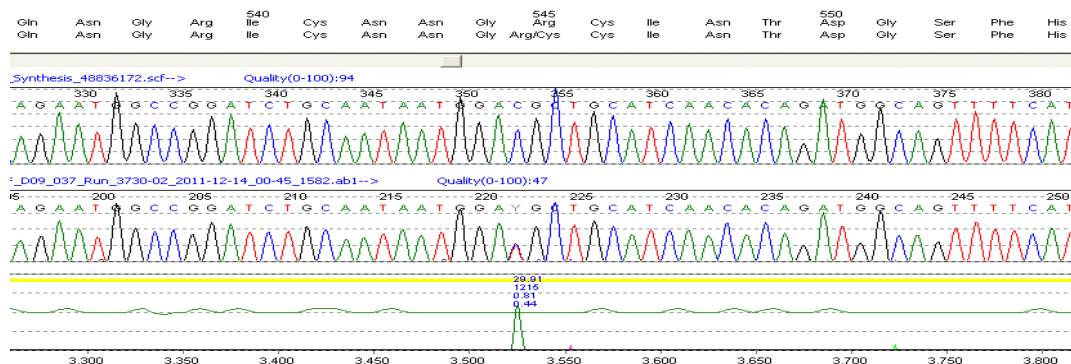
Cummings et al., 1998 JBJS

Arterial Tortuosity

- Tortuosity, elongation, and aneurysms of major arteries and the aorta
- Aortic stenosis, pulmonary artery or pulmonary valve
- Hypertelorism
- Hypermobile joints
- Arachnodactyly
- Scoliosis
- Hyperextensible skin
- Positive family history, sudden death in a close relative

Marfan Single Gene Assay

- 66 exons
- Mutation positive (~10% positivity rate)
 - Includes known pathogenic and suspected pathogenic
 - 56%: diagnosis based on clinical phenotype
 - 44%: suspected diagnosis of Marfan disease
- ~4% Variants of uncertain clinical significance
 - 64%: suspected diagnosis of Marfan
 - 37%: diagnosis based on clinical phenotype



Clinical Sensitivity of Gene Panel

- Aortopathy panel:
 - 17 genes
 - Each has clinical validity/utility separately
 - Clinical sensitivity: approximately 20% (doubled)
 - Internal data from Dr. P Bayrak-Toydemir

Looking towards the Future: Exome Diagnostic Yield

- Overall
 - 25%
 - N Engl J Med 2013; 369:1502-1511
- Severe Intellectual Disability: 16%
 - N Engl J Med 2012; 367:1921-1929
- Neurological diseases: 64%
 - [Brain](#). 2015 Feb;138(Pt 2):276-83.
- Retinal dystrophies: > 50%
 - Am J Ophthal online April 2015 [doi:10.1016/j.ajo.2015.04.026](https://doi.org/10.1016/j.ajo.2015.04.026)

Levels of Evidence

- Multiple models needed
 - Randomized control studies
 - Retrospective
 - Adaptive clinical trials
 - Diagnostic yield
 - Observational data
 - Linkage
 - Functional studies
 - Biological relationships/pathways
 - Current care vs molecular diagnostic models
 - Professional organization practice guidelines

Thanks to:

- AMP's Committees
 - Professional Relations
 - FEND working group
 - Clinical Practice
 - Economic Affairs
- ARUP Molecular Genetics/Genomics



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